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Maximum Match 100%
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Maximum DB seq length: 200000000
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Pred. No. is the number of results predicted by change to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SHIMMARIES

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"Function to Expert Themseleuses in Promobern.";

Submitted (JAN 1994) to the EMBG. Stubick, copy has been emBG. All H. C. (Lance All H. C.)
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EMBG AJ271297; CAC24721.1;
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of JHN Jord (TEMBELE), 12, Last septembre update)
FANCKEATIC RIBONUCLEASE PRECURSOR (FRACMENT).
                                                               N: VH.
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Heintema J.J.:
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Best Local Similarity 83.3%;
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Hest Local Similarity
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                                         bubalis."
                                                                                                                                                                                                                                                                                                                                                          Rubalus buhalis (Domestie Water buttalo).
Eukaryota; Metazod: Chordata: Traniata: Vertebrata: Euteleostomi;
Mammalia; Eutheria: Tetarti-da tyla; Ruminantia; Peccra; Hovoidea;
Hovidae: Bovinae: Bubalus.
                                                                                                                Vento M.T., Furia A.
                                                                                                                                                          Contalone E., Beintema J.J., Sisso M.P., Carsana A., Palmieri M.,
                                                                                                                                                                                                STRAIN- RIVER-TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01 007:2000 (TEMMRIOL. 15, Created)
01 007:2000 (TEMMRIOL. 15, List sept)
01-010-2000 (TEMMRIOL. 17, List and
2100:201 (TEMMRIOL. 17, List and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03.8 ZW60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hreukelman H.J., Warmels H.W., Beintema J.J.;
"Panereatie type fibonucleuses in Pronghorn.";
"Panereatie (yes) to be bed greenen/fibbl databases immortal action and communication and communica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dubois J.Y.F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antilocapia americana (Prosiborn).
Dakaryata, Mctacaa, Chandata. Yaniata, Vertebiata; Buteleostomi;
Mammalia: Butheria, Petarti daatyla, Bumimantia, Pecora, Bezoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      91 JUN 2001 (TERMELTEL 17, Created)
01-JUN 2001 (TERMELTEL 17, Cast annotation update)
01 JUN 2001 (TERMELTEL 17, Cast annotation update
Jabinitted (APK 2005) to the 3865;MemBask/DDBJ databases
                                                                  "Genomic sequence incoding general ribonuclease from Bubalus
                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                     NCBI_TaxID-89462;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q9MZ 45;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antilocapridão: Antilocapia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q9BDC2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 CAYKITQANKII I VACIDANI MVOGGGGASV RO
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163 | 18097 MW;
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83.38: Prof. No. 4.46-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2: Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             last sequence update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               S wro 141:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FRAIN-TYPE RIBONUCLEASE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 B31C7654A982A1CF CRC64.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      124 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 6: Length 163;
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Hest Local Similarity 83.3

Matches 25; concernation
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Best Local Similarity
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Ptam: PP00774: rnas6A.
Ptam: PP00774: rnas6A: 1.
PRINIS: PR00774: R160NUDIEASE.
ProDom: P1000535: RNAS6_Pc: 1.
SMART: SM00092: RNAS6_Pc: 1.
SMART: SM00092: RNAS6_Pc: 1.
PR05TIE: PS000127: RNASE_PANCREATIC: 1.
Endonuclease: Bydishase: Nuclease.
ACT_SITE 67
ACT_SITE 
      QRRPC2:
01-JUN-2001 (FEMPCH-), 17, Greated)
01-JUN-2001 (TEMPCH-), 17, Last sequence update)
01-JUN-2001 (TEMPCH-), 17, Last sequence update)
SEMINAL-TYPE RIBONUCLEASE (ERAUMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RUSSO N., de Higris M., Clardiello A., Di bonato A., D'Alessio G., "Baprossion in matematical cells, parification and characterization of recombinant buman par creatic ribonuclease."; pres fort 464-352-45 (1946) 1 SIMILARITY: TO THE PANCREATIC RIBONUCLEASE FAMILY. EMBL; $79281; AAB35096.1; -. HSSP; P00656; ISBN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-007 (IFEMBLE: 1. 01, Created) 01-01-07 (IFEMBLE: 1. 01, Lost sequence operate) 01-070-2001 (TrEMBLE: 1. 01, Lost annotation update) PANCREATIC RIFONUCLEASE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bukaryota, Metazoa, Chordata, Graniata, Vertebrata, Esterieostomi;
Mammalia: Butheria, Frimatos: Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                   Q9BEC2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE 95377432, CLEME  764928;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID-9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens (Human).
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Ptam: PF00074; rnaseA; ].
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EMBL; AF259584; AA-75.146.1; -.
                                                                                                                                                                                                                                                                                                                                     121 CAYRTSPKERHIIVACEGSPYVPVHFDASV 150
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                                                                                                                                                                                                                                                                                                                                                                                             1 CARKITCANEHIIVATEINEVADAHIDASA 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 CAYETTUMBELLIA BUNETVEVERLAGY 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRELIMINARY:
                                                                                                                                                                                   PRECIMINARY;
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WAse_Pe; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              89.7%; Score 134, DB 4, 79.7%; Pred. No. 1.18-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              4; Mismatches
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Fred. No. 4.4e 13,
0; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BY SIMILARITY.
B: FC2C981613ECEP92 CEC64:
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Interpres [PROUIAZZ] - ANASOA.
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J. Mol. Evol. 48: 600–68(1999).
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Obery Match 62.7% Grore 104; DB 6; Length 119; Best Lacal Smilarity 55.6% Grod. No. 3, Le-08; Matches 15; Conservative 6; Mismatches 6; Indels 0; Gaps 0;
```

search completed: March 14, 2092, 17:34:11 Job time: 652 sec

